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-Qu/cgn2 1/USPTO Bpool h/US09925867/runat 21032005 055412 1219/app query.fasta_1.391
-DB=Published Applications NA -QFMT=fastap -SUFFIX=rnpb -MINMATCH=0.1
-LOOPDLAO -LOOPDEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
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-MAXIEN=200000000 -USER=US09525867 @CGN 1 1 480 @runat 21032005 055412 1219
-NCPU=6 -ICPU=3 -NO MYAP -LARGEQUERY -NEG SCORE=9c -WAIT -DSPBLOCK=100
-LONGLOG -DEV_TIMEOŪT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5
-FGAPOP=6 -FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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     5544816 seqs,
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Ygapop 10.0 , Ygapext
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/cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
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/cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US11_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US60_NEW_PUB.seq:*
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2248.292 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Sequence 1, Applequence 1030, Applequence 5030, Applequence 5030, Applequence 5030, Applequence 149, Applequence 1662, Applequence 1662, Applequence 18631, Applequence 30088, equence 30088, equence 37889, applequence 27, Applequence 37, Applequence 32, A	%

ALIGNMENTS

RESULT 1
US-10-037-270-670
; Sequence 670, Application US/10037270
; Publication No. US20030104529A1
; GENERAL INFORMATION; Tom
; APPLICANT: Tang, Y. Tom
; APPLICANT: Liu, Chenghua
; APPLICANT: Asundi, Vinod APPLICANT: APPLICANT:
APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: Zhou, rang Ma, Yunqing Wang, Dunrui Asundi, Vinod Zhang, Jie Ren, Feiyan Chen, Rui-hong Zhao, Qing A. Wehrman, Tom Xue, Aidong J. Yang, Yonghong Wang, Jian-Rui Zhou, Ping

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APPLICANT: DIMBARS, Radoje T.

TITLE OF INVENTION: No. US20030104529A1el Nucleic Acids and FILE REFERENCE: 784C172B

CURRENT APPLICATION NUMBER: US/10/037,270

CURRENT APPLICATION NUMBER: 09/52,317

PRIOR APPLICATION NUMBER: 09/52,317

PRIOR APPLICATION NUMBER: 09/52,317

PRIOR FILING DATE: 2000-04-25

PRIOR FILING DATE: 2000-01-21

NUMBER OF SEQ ID NOS: 1104

SOFTMARE: 9t FL_genes Version 1.0

SEQ ID NO 670

LENGTH: 772

TYPE: DNA

ORGANISM: Homo sapiens

FEATURE:

**NAME/VEV. COO.**
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DB:
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; LOCATION: (24)..(665)
US-10-037-270-670
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IleLysArgGluArgArgLeuGlnIleTrpTyrArgArg 213
                                                                                   ProGlyCysProProThrAlaGluAlaLeuLeuTyrGlyIleLeuGlnLeuGlnArgLys 200
                                                                                                                                                                                CACTATTCCTACTCGGTGGGGGGGGGGGCGCGACGGATCGTGCCCGTGGACATCTACATC
                                                                                                                                                                                                                        HisTyrSerTyrSerValValArgGlyCysAspArgIleValProValAspIleTyrIle
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Matches:
Conservative:
Mismatches:
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Sequence 670, Application US/10117722

| Publication No. US20030219744A1
| GENERAL INFORMATION:
| APPLICANT: Tang, Y. Tom
| APPLICANT: Liu, Chenghua
| APPLICANT: Liu, Chenghua
| APPLICANT: Asundi, Vinod
| APPLICANT: Dranac, Radoje T.
| APPLICANT: Dranac, Radoje T.
| TITLE OF INVENTION: No. US20030219744A1el Nucleic Acids and
| TITLE OF INVENTION: Polypeptides
| FILE REFERENCE: 784CIPLECT
| CURRENT FILING DATE: 2002-04-04
| PRIOR APPLICATION NUMBER: US/10/117,722
| PRIOR APPLICATION NUMBER: 09/620,312
| PRIOR FILING DATE: 2000-01-9
| PRIOR APPLICATION NUMBER: 09/480,725
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; LOCATION: (24)..(665)
US-10-117-722-670
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US-10-117-722-670
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ORGANISM: Homo s
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No.:
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                                              ATCGTGGCCGGCACACTCACCAACAAGATGGCCCCAGCGCTTCGCAAGGTCTACGACCAG
                                                                                                                                                                                                         TyrAspMetAspArgPheGlyValValPheArgAlaSerProArgGlnSerAspValMet 120
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; SOFTWARE: PatentIn Ver. 2

; SEQ ID NO 625

; LENGTH: 2818

; TYPE: DNA

; ORGANISM: Homo sapiens

US-10-104-047-625
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; Publication No. US20030236392A1
; GENERAL INFORMATION:
   APPLICANT: HELIX RESEARCH INSTITUTE
   TITLE OF INVENTION: No. US20030236392A1e1 fu
; FILE REFERENCE: H1-A0105
; CURRENT APPLICATION UNMBER: US/10/104,047
; CURRENT FILING DATE: 2002-03-25
; PRIOR APPLICATION NUMBER:
; PRIOR APPLICATION NUMBER:
; PRIOR APPLICATION NUMBER:
; PRIOR PRIOR DATE:
; NUMBER OF SEQ ID NOS: 4096
; SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                 GAGTATGT
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            IleValAlaGlyThrLeuThrAsnLysMetAlaProAlaLeuA::gLysValTyrAspGln
                                                    TyrAspMetAspArgPheGlyValValPheArgAlaSerProArgGlnSerAspValMet
                                                                                                           CCCATGACCTTCGGCCTGGCCTGCCGCCGTGGAGATGATGCACATGGCAGCACCCCGC
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ATCGTGGCCGGCACACCAACAAGATGGCCCCAGCGCTTCGCAAGGTCTACGACCAG
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Sequence 914, Application US/10104047

| Publication No. US20030236392A1
| GENERAL INFORMATION:
| APPLICANT: HELIX RESEARCH INSTITUTE
| APPLICANT: HELIX RESEARCH INSTITUTE
| TITLE OF INVENTION: NO. US20030236392A1e1 full length cDNA
| FILE REFERENCE: H1-A0105
| CURRENT APPLICATION NUMBER: US/10/104,047
| CURRENT FILING DATE: 2002-03-25
| PRIOR APPLICATION NUMBER:
| PRIOR APPLICATION NUMBER:
| PRIOR FILING DATE: 2002-03-25
| VINDER OF SEQ ID NOS: 4096
| SOFTWARE: PATENTIN VET. 2.1
| SEQ ID NO 914
| LENGTH: 3089
| TYPE: DNA
| CONTINE NUMBER: US/10/104/1047
| LENGTH: 3089
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Best Local Similarity:
Query Match:
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                                                            SerThrGlnProAlaLeuProLysAlaArgAlaValAlaProLysProSerSerArgGly 60
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GluTyrValValAlAlALysLeuAspAspLeuValAsnTrpAlaArgArgSerSerLeuTrp
                                                                                                    ATCCGCCTCTGGGAAGCACCTGCGTGGCTGACGCCTCCTGTGCCCGTGTGTCTCTGTGCC
                                                                                                                                                           ACATCCCAGTCCCTGACCTCATGTGGGTCCAGGCCTCTGGCAGCGGCCGTGGGGGGCTCGC 351
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Mismatches:
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Sequence 262, Application US/10104047

Publication No. US20030236392A1

GENERAL INFORMATION:
APPLICANT: HELIX RESEARCH INSTITUTE
TITLE OF INVENTION: No. US20030236392A1el full length cDNA
CURRENT APPLICATION NUMBER: US/10/104,047

CURRENT FILING DATE: 2002-03-25

PRIOR APPLICATION NUMBER:
NUMBER OF SEQ ID NOS: 4096
SOFTWARE: PATENTIN VET. 2.1

SEQ ID NO 262
I SEQ ID NO 262
I SEQ ID NOS: 4096
ORGANISM: Homo sapiens
US-10-104-047-262
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Best Local Similarity:
Query Match:
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US-10-104-047-262
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Sequence 14502, Application US/10767701

Publication No. US20040172684A1

GENERAL INFORMATION:
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nuclecic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21(53535)B
CURRENT FILING DATE: 2004-01-29
NUMBER OF SEQ ID NOS: 63128
SEQ ID NO 14502
LENGTH: 1092
TYPE: DNA
ORGANISM: Sorghum bicolor
FEATURE:
OTHER INFORMATION: Clone ID: SORBI-28MAY03-CLUS4585_1
US-10-767-701-14502
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US-10-767-701-14502
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                pGlyProSerSerThrGlnProAlaLeuProLyBAlaArgAlaValAlaProLyBProSe
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 GGGCGCCTCCTCGACGTCCCCAGCG---CCATACGGGGGCGCGCCCCCCCCCGCCGGCGATGTC
                                                                             AlavalGlnAlaArgGly-Val----
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66.36%
66.23%
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Matches:
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Mismatches:
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APPLICANT: Zhou, Yihua

APPLICANT: Kovalic, David K.

APPLICANT: Screen, Steven E

APPLICANT: Tabaska, Jack E

APPLICANT: Tabaska, Jack E

APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

FILE REFERENCE: 38-21(53313)B

CURRENT APPLICATION NUMBER: US/10/425,114

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NO16284

LENGTH: 886

TYPE: nv"
                                                                                             US-09-525-867-1 (1-213)
                                                                                                                                               Query Match:
                                                                                                                                                            Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                              Score:
                                                                                                                                                                                                            Pred. No.:
                                                                                                                                                                                                                             Alignment Scores:
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                                                                                                                                                                                                                                                                                              FEATURE:
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ORGANISM: Zea mays
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AlaArgGly-Val-----
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                                  CCTCGTCCACGAGGTGAGGAAACCCTAGCGACCGACCATGGCGTTGCTCCCGCGCACCGC
                                                               ProGlyLeuArgGlyPheArgIleLeuGlyLeuArgSerSerValGlyProAlaValGln 26
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                                                                                                                               2.18e-85
737.50
75.70%
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                                                                                               x US-10-425-114-16284 (1-886)
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Matches:
Conservative:
Mismatches:
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Gaps:
 -----HisGlnSerValAlaThrAspGlyProSe
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
       US-09-525-867-1
                                                                                                Alignment Scores:
                                                                                                                      ; OTHER INFORMATION: Clone ID: US-10-425-114-6711
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US-10-425-114-6711
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                                                                                                                                                                   APPLICANT: Liu, Jingdong
APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT APPLICATION NUMBER: US/10/425,114
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 6711
LENGTH: 949
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Publication No. US20040034888A1
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                                                                                                                                                     TYPE: DNA
ORGANISM: Zea mays
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      (1-213)
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                          2.39e-85
737.50
75.70%
66.82%
65.97%
x US-10-425-114-6711 (1-949)
                                                                                                                           700576010_FLI
                                  Length:
Matches:
Conservative:
Mismatches:
Indels:
                     949
143
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RESULT 10
US-10-425-115-3877
US-10-425-115-3877
Sequence 3877, Application US/10425115
Publication No. US20040214272A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Clou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21 (53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
SEQ ID NO 3877
LENGTH: 1394
                                                                                                                Alignment Scores: Pred. No.:
                 Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                               US-10-425-115-3877
                                                                                                                                                                                                                                                                                                 LENGTH: 1394
TYPE: DNA
ORGANISM: Zea mays
                                                                                                                                                                                                                                    OTHER INFORMATION:
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             4.03e-85
737.50
75.70%
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                              Length:
Matches:
Conservative:
Mismatches:
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37

878

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995

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APPLICANT: LA ROSA, Thomas J.

APPLICANT: Kovalic, David K.

APPLICANT: Zhou, Yihua

ITILE OF INVENTION: Nucleic Acid Molecules and Oth

ITILE OF INVENTION: Plants

CURRENT APPLICATION NUMBER: US/10/425,115

CURRENT APPLICATION NUMBER: US/10/425,115

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 369326

SEQ ID NO 3875

LENGTH: 1840

TYPE: DNA

ORGANISM: Zea mays

FEATURE:
COTHER INFORMATION: Clone ID: MRT4577_103537C.1

US-10-425-115-3875
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DB:
Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 3875, Application US/10425115 Publication No. US20040214272A1 GENERAL INFORMATION:
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                       APPLICANT: Eroshkin, Alexey M.
APPLICANT: Zamudio, Carlos
TITLE OF INVENTION: IDENTIFICATION OF ESSENTIAL GEN
TITLE OF INVENTION: METHODS OF USE
FILE REFERENCE: 10182-021-999
CURRENT APPLICATION UMBER: US/10/320,797
CURRENT FILING DATE: 2002-12-16
PRIOR APPLICATION NUMBER: 60/341,261
PRIOR FILING DATE: 2001-12-17
NUMBER OF SEQ ID NOS: 3361
SOFTWARE: Patentin version 3.1
SOFTWARE: Patentin version 3.1
TYPE: DNA
TYPE: DNA
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US-10-320-797-2049
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ORGANISM: Cryptococcus neoformans
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Sequence 6216, Application US/10424599

Publication No. US20040031072A1

GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Cao Yongwei

TITLE OF INVENTION: Soy Nucleic Acid Molecul
TITLE OF INVENTION: Plants and Uses Thereof
FILE REFERENCE: 38-721(53223) B

CURRENT FILING DATE: 2003-04-28

NUMBER OF CONTROL

NUMBER: US/10/424,599
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US-10-424-599-6216/c
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NUMBER OF SEQ ID NOS: 285684
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RESULT 14
US-10-425-114-36399
Sequence 36399, Application US/10425114
Publication No. US20040034888A1
GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven B
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
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RESULT 15
US-10-424-599-6217/c
US-10-424-599-6217, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 36399
LENGTH: 863
TYPE: DNA
APPLICANT: LA ROSA Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
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ORGANISM: Glycine
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                                           cártactctractccgragtrcggggargrgacaggarrgrtccrgtrgacatatatgtr
                                                                                                         HisTyrSerTyrSerValValArgGlyCysAspArgIleValProValAspIleTyrIle
                                                                                                                                                   MetProGluProArgTyrValValSerMetGlySerCysAlaAsnGlyGlyGlyTyrTyr
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                                                                                                                                                                                                                                     TACGATCTCGACCGCTTCGGCATCATTTTCAGGCCCAGCCCTCGCCAGTCTGATTGCATG
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                                                                                                                                        ATGCCTGAGCCTAGATGGGTCGTCTCAATGGGAAGTTGTGCTAATGGAGGAGGATACTAC
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Search completed: March 22, 2005, 07:40:37 Job time: 582 secs

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Result
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-Q=/Cgn2 1/USFTO spool h/USO9525867/runat 21032005 055410 1201/app_query.fasta_1.391
-Q=/Cgn2 1/USFTO spool h/USO9525867/runat 21032005 055410 1201/app_query.fasta_1.391
-DB-ISBUEG Patents NA -QFMT=fastap -SUFFIX=rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=51ts -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000
-USER=USO9525867 @CGN 1 1 69 @FUNDAT = 21032005 055410 1201 -NCPU=6 -ICPU=3
-NO MMAP -LARGEQUERY -NEG SCORES=0 -WAIT -DSPELOCK=100 -LONGLOG
-DSV TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-DSV TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
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1: /cgn2_6/ptcdata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptcdata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptcdata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptcdata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptcdata/1/ina/Dackfiles1.seq:*
6: /cgn2_6/ptcdata/1/ina/Dackfiles1.seq:*
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3 US-09-103-840A-2

3 US-09-103-840A-1

US-09-103-840A-1

US-09-252-991A-14622

US-09-252-991A-15142

US-09-523-991A-15142

US-09-543-681A-43

US-09-596-002-30

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Sequence 14895, A
Sequence 2829, Appli
Sequence 2, Appli
Sequence 14622, A
Sequence 14622, A
Sequence 5374, Ap
Sequence 15142, A
Sequence 43, Appl
Sequence 30, Appli
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## ALIGNMENTS

US-09-620-312D-670

Sequence 670, Application US/09620312D

Patent No. 6569662

GENERAL INFORMATION:

APPLICANT: Tang, Y. Tom
APPLICANT: Tang, Y. Tom
APPLICANT: Asundi, Vinod
APPLICANT: Asundi, Vinod
APPLICANT: Chen, Rui-hong
APPLICANT: Chen, Rui-hong
APPLICANT: Chen, Rui-hong
APPLICANT: Webraman, Tom
APPLICANT: Wang, Jian-Rui
APPLICANT: Yang, Yonghong
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Jinwrui
APPLICANT: Wang, Dinurui
APPLICANT: Wang, Dinurui
APPLICANT: Wang, Dinurui
APPLICANT: John Tillinghast
APPLICANT: John Tillinghast
APPLICANT: John Tillingber
APPLICANT: John Tillingber
APPLICANT: John Tillingber
APPLICANT: John Tillingber
APPLICANT: OF INVENTION: No. 6569662el Nucleic Acids and
TITLE OF INVENTION: No. 6569662el Nucleic Acids and
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 78401P2B
CURRENT APPLICATION NUMBER: US/09/620,312D
CURRENT APPLICATION NUMBER: 09/552,317
PRIOR APPLICATION NUMBER: 09/552,317
PRIOR FILLING DATE: 2000-04-25
PRIOR FILLING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 1105
SOPTWARE: pt. FL genes Version 1.0
SEQ ID NO 670
LENGTH: 772
TYPE: DNA

sapiens

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i Sequence 1485, Application US/09270767
i Patent No. 6703491
i GENERAL INFORMATION:
i APPLICANT: Homburger et al.
i TITLE OF INVENTION: Nucleic acids and proteins of Dr.
i FILE REFERENCE: File Reference: 7326-094
i CURRENT APPLICATION UNUBER: US/09/270,767
i CURRENT ETLING DATE: 1999-03-17
i NUMBER OF SEQ ID NOS: 62517
i SOFTWARE: Patentin Ver. 2.0
i SEQ ID NO 14895
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; Sequence 2829, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANG
FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-13
; PRIOR APPLICATION UNMBER: US 60/074,725
; PRIOR APPLICATION UNMBER: US 60/074,725
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
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; SEQ ID NO 2829
LENGTH: 615
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SEQ ID NO 2
LENGTH: 4403765
TYPE: DNA
TYPE: DN
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; ORGANISM: Candida albicans
US-09-248-796A-2829
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                                                                                                                                                                                                                                                                                                   APPLICANT: FLEISCHMAN, Robert D.
APPLICANT: WHITE, Owen R.
APPLICANT: FRASER, Clair'S M.
APPLICANT: FRASER, Clair'S M.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
FULL REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
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7 AGAGTTATC 615
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Alignment Scores:

RESULT US-09- ; Seque ; pate ; papp ;	Qy dd	Qy VQ	Qy Db	DЬ	D Qy	Qy Db	ОУ	ОУ	Qy dd	B &	dd VQ	us-o	Pred. N Score: Percent Best Lo Query M DB:
13-840A ance 1, at No. 6 At No	195 3507939	175 3507879	155 3507822	135 3507762	115 3507702	96 3507642	76 3507582	56 3507534	36 3507516	3507495		9-525-867-	. No.: e: sent Similarity: Local Similarity: y Match:
pplicati 94328 MATION: LEISCHMP WHITE, C FRASER, VENTERNION: ENTION: CE: 2436 CE: 2436 ICATION	LeuGlnLeuGlnArgLys     :::    CTGAAGCTGCACGAAAAG	ProValAspIl             CCGGTCGACAT	AsnGlyGlyTyrTyrHis :::        TCGTCAGGTGGGATGTTCAAC	ArgLysValTy    ::::::   CGCCAGATCTA	ArgGlnSerAs       :::   CGGCAGGCAGA	MetAlaAlaPr     ACCGCGGGACC	ArgSerSerLe	ProSerSerAr	ThrAspGlyPr	GlyLeuArgSerSerValGly           ggATTGAGGTAGGGCGTGGGA	LeuSerAlaPr ::: ATTCATGCTCA	1 (1-213) ×	
US/091038  Robert D.  I R.  II C M.  SEQUENCE  A SEQUENCE  DOOD 7.00  OBER: US/0  1998-06-24	nArgLysIleLys      ::: CGAAAAGATTCAG	ProValAspIleTyrIleProGlyCys	YTyrTyrHisT   :::::  GATGTTCAAC-	rAspGlnMetE           TGACCAGATGO	PValMetIle\   :::        TCTGATGATCC	OArgTyrAsp\     :::     AAGGTTTGAC <i>I</i>	uTrpProMet]          GTGGCCGGCA/	gGlyGluTyr\     -GGCGGGATCO	oSerSerThro	rSerValGlyF	ProGlyLeuArgGlyPheArgIl    :::       cacggrgTTCGTGGCCTACGCGT	US-09-103-	1.99e-37 449.00 60.91% 41.82% 40.16%
40A S FOR STRAIN S 9/103,840A	IleLys       ;;   column	ProValAspIleTyrIleProGlyCysProProThrAlaGluAlaLeuLeuTyrGlyIl              :::             CGGTCGACATCTACCTACCCGGCTGCCGCGCGCGGAGATGCTGCTGCACGCAAT	· <	ArgLysValTyrAspGlnMetProGluProArgTyrValValSerMetGlySerCysAla     :::::              -    :::::	ArgGlnSerAspValMetIleValAlaGlyThrLeuThrAsnLysMetAlaProAlaLeu       :::  :::                	MetAlaAlaProArgTyrAspMetAspArgPheGlyValValPheArgAlaSerPro	ArgSerSerLeuTrpProMetThrPheGlyLeuAlaCy8Cy8AlaValGluMetMetHi8 :	rSerArgGlyGluTyrValValAlaLaLysLeuAspAspLeuValAsnTrpAlaArg 	ThrAspGlyProSerSerThrGlnProAlaLeuPro        CTGGAAGAACAGCTGCCC	rSerValGlyProAlaValGlnAlaArgGlyValHisGlnSerValAla        GGGCGTGGGA	LeuSerAlaProGlyLeuArgGlyPheArgIle	840A-2 (1-	Length: Matches: Conservat Mismatche Indels: Gaps:
Analysis	TTAGGTATCA	ProProThrAlaGluAlaLeuLeuTyrGlyIle	rSerTyrSerValValArgGlyCyeAspArgIleVal 	TyrValVals: :::   :::: TGGGTTCTGG	:LeuThrAsnL; :::::: ;GTCAGCCAGA	Glyvalval-    ::: GGGATGGAACO	Alacyscysa            GCGTGCTGTG	LeuAspAspLo :::::: :GTCGAGAAGG	ProLysAlaA      CCC	AlaArgGlyVa	TGTGTGGCGC	4403765)	4403 92 1tive: 42 1e8: 38
IN MYCOBACTERIUM	ArgGluArgArgLeu          xccgggaacgcgctatc	laLeuLeuTyr       :::  GCTGCTGCAC	LYCYBABDATG          3CGTGGATCAT	arMetGlySer ::      CATGGGTGTG	/sMetAlaPro           \GATGGCGCCG	PheArgAla     	laValGluMet   :::       GATCGAGATG	euValAenTrp ::: rGGCGGGCTAT	roLysAlaArgAlaValAlaProLys    CC	lHisGinSer	GCGGGGGCCT		1765
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Query Match: DB:

Score:

Alignment Scores:

H37Rv

No.:

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Sequence 14622, Application US/09252991A

Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: MATC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQ
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
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Best Local Similarity:
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Sequence 5374, Application US/09489039A
Patent No. 6610836
GENERAL INFORMATION:
APPLICANT: GATY Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: ENBUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
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CURRENT APPLICATION NUMBER: US/09/489,039A; CURRENT FILING DATE: 2000-01-27; PRIOR APPLICATION NUMBER: US 60/117,747; PRIOR FILING DATE: 1999-01-29; NUMBER OF SEQ ID NOS: 14342; SEQ ID NO 5374; LENGTH: 690
TYPE: DNA
ORGANISM: Klebsiella pneumoniae
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                 SEQ ID NO 15142
LENGTH: 696
TYPE: DNA
                                                                                                                                                                                                                                                                     Sequence 15142, Application US/09252991A Patent No. 6551795
                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                             CURRENT APPLICATION NUMBER: US/09/252,991A CURRENT FILING DATE: 1999-02-18 PRIOR APPLICATION NUMBER: US 60/074,788 PRIOR FILING DATE: 1998-02-18 PRIOR APPLICATION NUMBER: US 60/094,190 PRIOR FILING DATE: 1998-07-27 NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                 APPLICANT: MARC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
ORGANISM: Pseudomonas aeruginosa
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                                                                                                ; SEQ ID NO 43
; LENGTH: 735
; TYPE: DNA
; ORGANISM: Proteus m
US-09-543-681A-43
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Best Local Similarity:
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US-09-543-681A-43
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APPLICANT: CARY BRETON
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABIL
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.1002-001
CURRENT PELLICATION NUMBER: US/09/543,681A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR FILING DATE: 1999-04-09
NUMBER OF SEQ ID NOS: 8344
                                                                                                                                                                                                                                                                                                                                                        Sequence 43, Application Patent No. 6605709
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APPLICANT: Lagace, ROUGE,
APPLICANT: Patterson, Chandra
APPLICANT: Berg, Kim, L.
TITLE OF INVENTION: NUCLEOTIDE SEQUENCES OF
FILE REFERENCE: PM-0008-4 US
CURRENT APPLICATION NUMBER: US/09/596,002
CURRENT FILING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: 60/140,121
PRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS:
NUMBER OF SEQ ID NOS:
SOFTWARE: PERL Program
    US-09-525-867-1 (1-213)
                                   Query Match:
DB:
                                                      Percent Similarity:
Best Local Similarity:
                                                                                                                             ; FEATURE:
; NAME/KEY: misc_feature
; OTHER INFORWATION: Incyte template
; PUBLICATION INFORWATION:
US-09-596-002-30
                                                                                                                                                                                                  ; SOFTWARE: PERL Program
; SEQ TID NO 30
; LENGTH: 58909
; TYPE: DNA
; TYPE: DNA
; OFFERMISM: M. catarrhalis
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US-09-596-002-30
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                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 30, Application US/09596002
Patent No. 6632636
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Sequence 1, Application US/09790988

Patent No. 6632935

GENERAL INFORMATION:

APPLICANT: SHIGENOBL, SHUJI

APPLICANT: WATANABE, HIDEMI

APPLICANT: WATANABE, HIDEMI

APPLICANT: HATTORI, MASAHIRA

APPLICANT: HATTORI, MOSHITVIKI

ITITLE OF INVENTION: GENOME DINA OF BACTERIAL SYMBIONT OF APH

CURRENT APPLICATION NUMBER: US/09/790,988

CURRENT APPLICATION NUMBER: 2001-02-23

CURRENT APPLICATION NUMBER: US/09/790,988
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                                                            US-09-525-867-1 (1-213) x US-09-790-988-1 (1-640681)
                                                                                                                  Query Match:
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Best Local Similarity:
                                                                                                                                                                                             Alignment Scores:
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PRIOR FILING DATE: 2000-04-07
NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Buchnera
                                                                                                                                                                                                                                                                        LENGTH: 640681
                                                                                                                                                                                No.:
165027 TTACATAAATTAGTTAATTGGGGTCGAAAAAATTCTTTATGGCCTTATAATTTTGGTCTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AlaProAlaLeuArgLysValTyrAspGlnMetProGluProArgTyrValValSerMet 150
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428.00
72.41%
55.86%
38.28%
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Mismatches:
Indels:
Gaps:
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Matches:
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Percent Similarity:
Best Local Similarity:
Query Match:
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APPLICANT: GATY L. Breton et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING '
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2005-001

CURRENT APPLICATION NUMBER: US/09/540,236

CURRENT FILING DATE: 2000-04-04

NUMBER OF SEQ ID NOS: 3840

SEQ ID NO 467

LENGTH: 708

TYPE: DNA

ORGANISM: M.catarrhalis
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GENERAL INFORMATION:
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                                                                                                                                                                        CCGAGGTG-GATAAAAATGTTTTTTTGGGCAGATTGTCCGATTTGACGCACAACCTTGCC
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 ArgAlaSerProArgGlnSerAspValMetIleValAlaGlyThrLeuThrAsnLysMet 130
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                                                                   GluMetMetHiBMetAlaAlaProArgTyrAspMetAspArgPheGlyVal---ValPhe 110
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                                    GAGTATGCCACGACCTTGACAGGCGTGCATGATTTGTCTCGTTTTTGGGGCCGGAGGTCATT
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US-09-328-352-581
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 8252
SEQ ID NO 581
LENGTH: 684
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GENERAL INFORMATION:
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                                                    ATGTACGACATC---TATTCAGTAGTACAAGGTGTAGACAAAATTATTCCTGTTGACGTG
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                                                                                                                                                                                                                                                                                             ArgTyrAspMetAspArgPheGlyVal----ValPheArgAlaSerProArgGlnSerAsp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GlySerCysAlaAsmGlyGlyTyrTyrHisTyrSerTyrSerValValArgGlyCys 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AlaProAlaLeuArgLysValTyrAspGlnMetProGluProArgTyrValValSerMet 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TyrTyrHisTyrSerTyrSerValValArgGlyCysAspArgIleValProValAspIle 178
                                                                                                                       GAACAAATGTTAGAACCTAAATGGGTTATTTCAATGGGTGCTTGTGCAAACTCTGGAGGT
                                                                                                                                           TIGATGATTGTTGCAGGTACCTGCTTTGTAAAAATGGCTCCGGTTATCCAACGTTTGTAT
                                                                                                                                                                                                                  ValMetIleValAlaGlyThrLeuThrAsnLysMetAlaProAlaLeuArgLysValTyr 138
                                                                                                                                                                                                                                                                                                                                                                   TrpProMetThrPheGlyLeuAlaCysCysAlaValGluMetMetHisMetAlaAlaPro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GATAAAATCTTGCCTGTGGATGTGTATGTTCCGGGGTGTCCGCCACGCCCCGAGGCTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCACCTGTGATTCTCAGGCTGTATGAACAAATGCTAGAACCCAAGTGGGTCATCTCCATG
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Gaps:
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Matches:
Conservative:
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US-09-525-867-1 (1-213) x US-09-020-956-32 (1-789)
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                         Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                         ; MOLECULE TYPE: US-09-020-956-32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME:
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 2101
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEPAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 14
US-09-020-956-32/c
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Patent No. 6261562
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APPLICANT: Xu, Jiangchun
APPLICANT: Dillin, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF
NUMBER OF SEQUENCES: 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: YOLV;

COMPUTER READBLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

""""TOATTON NUMBER: US/09/020,956
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
TYPE: n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: 09-FEI CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
             534 CTGNTGNGCCCGGGAGGTGATGNANATNGCAGCAACCCCGNNAAGACATGGNCCGGTTTT 475
                                                                                         587
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                                      87 aCysCysAlaValGluMetMetHisMetAlaAla-ProArgTyrAspMetAspArgPhe- 106
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                                                                                                                                                                                                                                   36 ThrAspGlyProSerSerThrGlnProAlaLeuProLysAlaArgAlaValAlaProLys
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                                                                          -----AAACTGGGNCCCCCGGAGNTTTTTTGNGGGCCNCAAGAACTTTTGGGTTGNC
                                                                                                               gArgSerSerLeuTrpPro---
                                                                                                                                        CCCAGNAANCGGGGGGGGATTTGGGGGGCCAAGGGGGGATGACTTTGTTG-------
                                                                                                                                                                       ProSerSerArgGlyGluTyrValValAlaLaLysLeuAspAsp-LeuValAsnTrpAlaAr 75
                                                                                                                                                                                                       ACCGGGGGCCCGAAGGGGGANCCANCTTGCCNNCCAAAGNNAGAGGCGTGGGTTCCCCAAA
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          Percent Similarity:
Best Local Similarity:
Query Match:
DB:
                                                                        Score:
                                                                                    Alignment Scores: Pred. No.:
                                                                                                                                   ; MOLECULE TYPE: US-09-030-607-32
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                     FILING DATE: 25-FEB-1998
CLASSIFICATION:
ATTORNEY AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 2101
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Ku, Jiangchun C. APPLICANT: Dillon, Davin C. TITLE OF INVENTION: COMPOUND: NUMBER OF SEQUENCES: 224
CORRESPONDENCE ADDRESS:
                                                                                                                                                           LENGTH: 789 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/030,60
                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              146 rValValSerMetGlySerCysAlaAsnGlyGlyGlyTyrTyrHisTyrSerTyrSerVa
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5. 6262245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          126 uThrAsnLysMetAlaProAlaLeuArgLysValTyrAspGlnMetProGluProArgTy 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           107
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                                                                                                                                                                                                                                                         (206) 682-6031
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                              Length:
Matches:
Conservative:
Mismatches:
   Indels:
Gaps:
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	Db 177 GCTGCANATTTGGTACCGCAGG 156  Search completed: March 22, 2005, 07:50:56  Job time : 5030 Becs	Db 177 Search comple
TGCAGAGGAANATCAAGCGGGAGG		
REPILETYTILEPTOGLYCYSPTOP:	1ValargGlyCypasparglleValProValAspIleTyrIleProGlyCypProProTh	
31yG1yTyrTyrHisTyrSerTyrSe              GNGGGTAANACCACTATTCCTAATN	rValValSerMetGlySerCysAlaAsnGlyGlyGlyTyrTyrHisTyrSerTyrSerVa	Oy 146 Db 356
/alTyrAspGlnMetProGluProAr 	uThrAsnLysMetAlaProAlaLeuArgLysValTyrAspGlnMetProGluProArgTy 146	Qy 126 Db 415
erAspValMetIleValAlaGlyTh:                  CTGA-GTNATGATTGTGGCNGGNANJ	-GlyValValPheArgAlaSerProArgGlnSerAspValMetIleValAlaGlyThrLe	Qy 107 Db 474
lla-ProargTyraspMetaspargP 	aCysCysAlaValGluMetMetHisMetAlaAla-ProArgTyrAspMetAspArgPhe- 	Qy 87 Db 534
TGNGGGCCNCAAGAACTTTTGGGTT	gArgSerSerLeuTrpPro	Qy 75 Db 587
/sLeuAspAsp-LeuValAsnTrpAla                       : : : \GGGGGATGACTTTGTTG	ProSerSerArgGlyGluTyrValValAlaLysLeuAspAsp-LeuV&lAsnTrpAlaAr	Qy 56 Db 636
uProLysAlaArgAlaValAlaProl       	ThraspGlyProSerSerThrGlnProAlaLeuProLysAlaArgAlaValAlaProLys	Oy 36 Db 696
.789)	US-09-525-867-1 (1-213) x US-09-030-607-32 (1-789)	US-09-525-867

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Searched:
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Perfect score:
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Published Applications NA:*

1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
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12: /cgn2_6/ptodata/2/pubpna/US09C_PUBCOMB.seq:*
13: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*
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21: /cgn2_6/ptodata/2/pubpna/US10R_PUBCOMB.seq:*
22: /cgn2_6/ptodata/2/pubpna/US10R_PUBCOMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

000000 110987654221	Result
7500.8 569.2 427.2 384.6 384.6 384.6	Score
991. 991. 991. 446. 466.7 777.	% Query Match Length DB
772 2818 3050 3089 789 789 789 789 789	Length
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Sequence 4822, Ap	20.5		Sequence 6030, Ap	Sequence 1030, Ap	Sequence 2030, Ap	e 1,	4866	316,	273,	715,	Sequence 7030, Ap	Sequence 1, Appli	198			6216, A	6217, A	3877,	Sequence 6711, Ap		e 3875,	O	3639	9249		55,	32,	32,	e 32,	Sequence 32, Appl	e 32	w	Sequence 32, Appl

## ALIGNMENTS

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FITLE   FILE   CURRED CURRED PRIOR	APPLI APPLI APPLI APPLI	APPI APPI APPI APPI APPI APPI	RESULT 1  US-10-037-270-670  Sequence 670, Application US/10  Publication No. US20030104529A1  GENERAL INFORMATION:  APPLICANT: Tang, Y. Tom  APPLICANT: Lau, Chenghua  APPLICANT: Asundi, Vinod  APPLICANT: Chen, Rui-hong  APPLICANT: Chen, Rui-hong
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INVENTION: INTERENCE: 784CII RRENCE: 784CII RRENCE: 784CII RRENCE: 784CII RRENCE: 200 LICATION NUM LICATION N	Wang, Tilli Tilli Drman	Zhao, Wehrm Xue, Yang, Yang, Wang, Zhou, Ma, Y	670 No. Dpplica No. US200 No. US200 RMATION: Tang, Y. Liu, Che Asundi, Zhang, Zhang, Chen, Re
CATTON: 784 ICATION NG DATE ATION N DATE: DATE: DATE: DATE: CFL_ger	lin lin		Applico US2.  MATION MATION Cliu, Cliu, Cliu, Clandiang, Franc, Franc, Chen, I
Y ACIP ACIP N NU E: NUMB 200 200 NUMB 200 NUMB NUMB NUMB NUMB NUMB	Zhiwei ghast, ghast, c, Rad	zhao, Qing A. wehrman, Tom Xue, Aidong J Yang, Yonghon Wang, Jian-Ru Zhou, Ping Zhou, Ping Zhou, Ping Zhou, Ping Zhou, Ping	70 Application o. US20030104 MATION: ang, Y. Tom any, Chenghua Liu, Chenghua Asundi, Vinod Zhang, Jie Ren, Feiyan Chen, Rui-hon
FOLYPEPT CIP2B I NUMBER: U : 2002-01 UMMBER: 09/ 2000-04-25 UMMBER: 09/ 2000-01-21 SS: 1104 les Version	Wang, Zhiwei Wang, Zhiwei Tillinghast, John Drmanac, Radoje T ENTION: No. US200	nan, Tom Aidong J. Yonghong Yian-Rui Ping Yunging	cation U 100301045 1. Tom henghua henyan Vinod Jie Viyan Rui-hong
OF INVENTION: POLYPEDICIDES REFERENCE: 784CIPES NT APPLICATION NUMBER: US/10/037,270 NT FILING DATE: 2002-01-04 APPLICATION NUMBER: 09/552,317 FILING DATE: 2000-04-25 APPLICATION NUMBER: 09/488,725 FILING DATE: 2000-01-21 R OF SEQ ID NOS: 1104 ARE: pt_FL_genes Version 1.0 NO 670	Wang, Zhiwei Tillinghast, John Drmanac, Radoje T. INVENTIN: No. US20030104529A1el		270-670 670, Application US/10037270 ion No. US20030104529A1 INFORMATION: INT: Tang, Y. Tom NT: Liu, Chenghua NT: Asundi, Vinod NT: Asundi, Vinod NT: Ren, Feiyan NT: Ren, Feiyan NT: Chen, Rui-hong
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RESULT 2
US-10-117-722-670
IS-quence 670, Application US/10117722
Publication No. US20030219744A1
GENERAL INFORMATION:
APPLICANT: Tang, Y. Tom
APPLICANT: Liu, Chenghua
APPLICANT: Asundi, Vinod
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: ORGANISM: Homo sapiens
: FEATURE: Homo sapiens
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APPLICANT: Zhang, Jie
APPLICANT: Drmanac, Radoje T.
ITILE OF INVENTION: No. US20030219744A1el Nucleic Aci
ITILE OF INVENTION: Polypeptides
FILE REFERENCE: 784CIPZBCIP
CURRENT APPLICATION NUMBER: US/10/117,722
FRIOR APPLICATION NUMBER: 09/620,312
FRIOR APPLICATION NUMBER: 09/620,312
FRIOR APPLICATION NUMBER: 09/52,317
FRIOR FILING DATE: 2000-07-19
FRIOR APPLICATION NUMBER: 09/552,317
FRIOR FILING DATE: 2000-04-25
FRIOR FILING DATE: 2000-01-21
FRIOR FILI
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; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (24)..(665)
US-10-117-722-670
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99.1%; Pred. No. 3.7e-199;
tive 0; Mismatches 7;
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CCGCCGCCGCCGCCGGAGCCTGTCGCCGTCCTGTCCCCAGCCTGCTTGTGTCCCCGTG

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APPLICANT: HELIX RESEARCH INSTITUTE
ITITE OF INVENTION: NO. US20030236392A1el full length cDNA
FILE REFERENCE: 11-A0105
CURRENT APPLICATION NUMBER: US/10/104,047
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER:
PRIOR APPLICATION NUMBER:
PRIOR APPLICATION NUMBER:
PRIOR FILING DATE:
NUMBER OF SEQ ID NOS: 4096
SOFTWARE: PATCHING Ver. 2.1
SEQ ID NO 625
LENGTH: 2818
TYPE: DNA
ORGANISM: Homo sapiens
US-10-104-047-625
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Best Local Simi
Matches 577;
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b. US20030236392A1
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Pred. No. 1.4e-148;
0; Mismatches 13;
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Publication No. US20030236392A1

GENERAL INFORMATION:
APPLICANT: HELIX RESEARCH INSTITUTE
ITITLE OF INVENTION: NO. US20030236392A1e1 fu

FILE REFERENCE: H1-A0105
CURRENT APPLICATION NUMBER: US/10/104,047
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER:
                                                                                                                                                                  RESULT 5
US-10-104-047-914
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-104-047-262
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Sequence 914, Application US/10104047
Publication No. US20030236392A1
GENERAL INFORMATION:
APPLICANT: HELIX RESEARCH INSTITUTE
TITLE OF INVENTION: No. US20030236392A1e1 full length
FILE REFERENCE: H1-A0105
CURRENT APPLICATION NUMBER: US/10/104,047
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 4096
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 262
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Best Local Similarity
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Pred. No. 8.4e-127;
0; Mismatches 15;
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Sequence 32, Application US/09759143

Fatent No. US202022248A1

FAPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Henderson, Robert A.
APPLICANT: Retter, Marc W.
APPLICANT: Retter, Marc W.
APPLICANT: Bob, Craig H.
APPLICANT: Day, Craig H.
APPLICANT: Loaver, Carter, Darrick
APPLICANT: Li, Samuel
APPLICANT: Li, Samuel
APPLICANT: Keeky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHOD
FITLE REFERENCE: 210121.427C23
CURRENT APPLICATION NUMBER: US/09/759,143

CURRENT FILING DATE: 2001-01-12
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US-09-759-143-32/c
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SOFTWARE: PatentIn v
SEQ ID NO 914
LENGTH: 3089
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ORGANISM: Homo sapiens
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Pred. No. 4.9e-109;
0; Mismatches 13;
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SEQ ID NO 32
LENGTH: 789
TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(789)
OTHER INFORMATION: n = A
US-09-759-143-32
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Best Local Similarity 75.9%;
Matches 527; Conservative
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                                              GCCGCCGGCCGGAGCCTGTCGCCGTCCTGTCCCCAGCCTGTCTTGTCTCCCGTGAGGTTGTC
                                                                            AGAGGAANATCAAGCGGGAGCGGAGGCTGCANATTTGGTACCGCAGGTAGGGCCGCCCCCC
                                                                                        AGAGGAAGATCAAGCGGGAGCGGAGGCTGCAGATCTGGTACCGCAGGTAGCGCCGCCGCC
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Pred. No. 3.4e-97;
0; Mismatches 154;
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Sequence 32, Application US/09780669
Patent No. US20020051977A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan L.

RESULT 7 US-09-780-669-32/c

Kalos, Michael

Yugui

Robert

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APPLICANT: Hural, John
APPLICANT: McNeill, Patricia D.
APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C24
CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 943
SOFTWARE: FASUSEQ for Windows Version 3.0
SEQ ID NO 32
SEQ ID NO 32
LENGTH: 789
TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
US-09-780-669-32
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Best Local Similarity 75.5
Matches 527; Conservative
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APPLICANT:
APPLICANT:
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                                                                  GCTACTACCACTATTCCTACTCGGTGGTGAGGGGGCTGCGACCGCATCGTGCCCGTGGACA
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       TCTACATCCCAGGCTGCCCACCTACGGCCGAGGCCCTGCTCTACGGCATCCTGCAGCTGC
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Hepler, William
Hural, John
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Vedvick, Thomas S.
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Pred. No. 3.4e-97;
0; Mismatches 154;
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US-09-030-606-32/c
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TELEPHONE: (206) 682-6031
INFORMATION FOR SEQ ID NO: 32
SEQUENCE CHARACTERISTICS:
LENGTH: 789 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 527; Conserv
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APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
TITLE OF INVENTION: COMPOUNDS
NUMBER OF SEQUENCES: 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/030
PILING DATE: 25-FEB-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REGISTRATION NUMBER: 2101
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1000: 122-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 98104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
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559
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GGGCCNCAAGAACTTTTGGGTTGNCCTGNTGNGCCCGGGAGGTGATGNANATNGCAGCAA
                   G----CCCATGACCTTCGGCCTGGCCTGCGCCCGTGGAGATGATGCACATGGCAGC-A
                                                                 GGATTTGGGGGCCAAGGGGGATGACTTTGTTGAAACTGGGNCCCCCCGGAGNTTTTTTGN
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75.9%;
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                                                                                                                                                                                                                                                                       Score 384.6; DB 9;
Pred. No. 3.4e-97;
0; Mismatches 154;
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; APPLICANT: Xu, Jiangchun
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND ITILE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
; FILE REFERENCE: 210121.534C1
; CURRENT APPLICATION NUMBER: US/09/822,827
; CURRENT FALING DATE: 2001-03-28
; NUMBER OF SEQ ID NOS: 982
; SOFTWARE: FASTSEQ for Windows Version 3.0
; SEQ ID NO 32
; SEQ ID NO 32
; SEQ ID NO 30
; SEQ ID NO 30
; SEQ ID NO 30
; AND TYPE: DNA
; ORGANISM: Homo sapien
; PATURE:
; NAME/KEY: misc feature
; COCATION: (1)...(789)
; OTHER INFORMATION: n = A,T,C or G
US-09-822-827-32
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; Sequence 32, Application US/09822827

; Patent No. US20020081680A1
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Best Local Similarity 75.9
Matches 527; Conservative
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APPLICANT: Xu, Jian
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Pred. No. 3.4e-97;
0; Mismatches 154
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US-09-113-453-32/c

| Sequence 32, Application US/09115453B |
| Patent No. US20020090372A1 |
| GENERAL INFORMATION: |
| APPLICANT: Xu, Jiangchun |
| APPLICANT: Dillon, Davin C. |
| TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF |
| TITLE OF INVENTION: METHODS FOR THEIR USE |
| FILE REFERENCE: 210121.427C4 |
| CURRENT APPLICATION NUMBER: US/09/115,453B |
| CURRENT FILING DATE: 1998-07-14 |
| NUMBER OF SEQ ID NOS: 228 |
| SOFTWARE: FastSEQ for Windows Version 3.0 |
| SEQ ID NO 32 |
| TENTET OF THE SECON NUMBER: VERSION 3.0 |
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TYPE: DNA
ORGANISM: Homo sapien
FEATURE: misc_feature
LOCATION: (1)...(789)
OTHER INFORMATION: n = A
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                       CACCCAGCCTGCCCTGCCAAAGGCCAGAGCCGTGGCTCCCAAACCCAGCAGCAGCCGGGCCGA 260
                                                                                               GGGGGATGGTGNNNAGGANGGNGTTTTCNTTAGAGNNGGGCCCCACCGGGGGCCCCGAAGGG
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Pred. No. 3.4e-97;
0; Mismatches 154;
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 GGGGGATGGTGNNNAGGANGGNGTTTTCNTTAGAGNNGGGCCCACCGGGGGGCCCGAAGGG
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                                                      Conservative
                                                                46.7%;
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                                                   Score 384.6; DB 9;
Pred. No. 3.4e-97;
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Sequence 32, Application US/098;
Publication No. US20020192763A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
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         Hural, John
McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Ca
Foy, Teresa
                                                                                                                                   Day, Craig H. Vedvick, Thomas Carter, Darrick Li, Samuel X.
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                                                                            Skeiky, Yasir A.W.
Hepler, William T.
Henderson, Robert A.
                                                                                                                                                                                      Kalos, Michael D.
Retter, Marc W.
Stolk, John A.
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Mitcham, Jennifer L.
                                                                                                                      Wang, Aijun
Fanger, Gary R.
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; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
; FILE REFERENCE: 210121.534C2
; CURRENT APPLICATION NUMBER: US/09/895,793
; CURRENT FILING DATE: 2001-06-29
; NUMBER OF SEQ ID NOS: 982
; SOFTWARE: FeatSEQ for Windows Version 3.0
; SEQ ID NO 32
; SEQ ID NO 32
; LENGTH: 789
; TYPE: DNA
; ORGANISM: Homo sapien
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(789)
; OTHER INFORMATION: n = A,T,C or G
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Sequence 32,
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          Application US/09895814
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Pred. No. 3.4e-97;
0; Mismatches 154;
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; OTHER INFORMATION: n = A,T,C or US-09-895-814-32
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Best Local Similarity
Matches 527; Conserv
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CURRENT FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 990
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
TYPE: DNA
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GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C26
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                                                                               A-GTNATGATTGTGGCNGGNANANTAACCAAAAAGAGGGCCCCAGNG
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Hepler, William T.
Henderson, Robert A.
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Pred. No. 3.4e-97;
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APPLICANT: Madicaline JO
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C27
CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT PAPLICATION OF SEQ 1D NOS: 1011
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
FEATURE:
FEATURE
VAMME/KEY: misc_feature
LOCATION: 65, 68, 92, 103, 171, 195, 219, 238, 240, 260, 288, 301, LOCATION: 317, 324, 327, 336, 345, 358, 395, 417, 419, 421, 424, 436, LOCATION: 449, 468, 470, 483, 493, 494, 507, 510, 512, 528, 531, 536, LOCATION: 554, 560, 568, 579, 628, 631, 656, 657, 664, 665, 672
VAMME/KEY: misc_feature
LOCATION: 745, 749, 753, 757, 759, 773, 787
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US-10-012-896-32/c
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US-10-012-896-32
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McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
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Hepler, William T.
Henderson, Robert A.
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Vedvick, Thomas S.
Carter, Darrick
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Harlocker, Susan L.
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Retter, Marc W.
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be, Yoshihiro
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Pred. No. 3.4e-97;
0; Mismatches 154;
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US-10-010-940-32/c

US-10-010-940-32/c

Sequence 32, Application US/10010940

Publication No. US20030088062A1

GENERAL INFORMATION:

APPLICANT: XL, Jiangchun

APPLICANT: Mitcham, Jennifer L.

APPLICANT: Mitcham, Jennifer L.

APPLICANT: Jiang Yuqui

APPLICANT: Jiang Yuqui

APPLICANT: Reed, Steven G.

APPLICANT: Ralos, Michael

APPLICANT: Fanger, Gary

APPLICANT: Retter, Mark

APPLICANT: Solk, John

APPLICANT: Day, Craig

ITILE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND

ITILE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
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CURRENT APPLICATION NUMBER: US/10/010,940
CURRENT FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
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               AATAAACCTGCCCTCGGGCAAAAAAAAAAAAAAAAAA 824
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Search co Job time

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Perfect score:
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        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/EB_COMB.seq:*

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Listing first 45 summaries
     Copyright
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US-09-620-312D-670
US-09-030-607-32
US-09-030-607-32
US-09-3313-32
US-09-352-616A-32
US-09-159-812-32
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Sequence 670, App
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RESULT 2
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Patent No.
STREET: 6300 COlumbia Center, 701 F
STREET: 6300 COlumbia Center, 701 F
STATE: WA
COUNTRY: USA
ZIP: 98104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Ve.
APPLICATION UNMER: US/09/020,956
FILING DATE: 09-FEB-1998
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APPLICANT: Xu, Jiangchun
APPLICANT: Dillin, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR
NUMBER OF SEQUENCES: 178
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ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center,
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INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
LENGTH: 789 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
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ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 2101
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                            GCCGCCGCCGGAGCCTGTCGCCGTGCCTGTCCCCAGCCTGCTTGTGTCCCGTGAGGTTGTC
AGAGGAANATCAAGCGGGAGCGGAGGCTGCANATTTGGTACCGCAGGTAGGGCCGCCGCC
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75.9%;
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Pred. No. 4.2e-80;
0; Mismatches 154;
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RESULT 3 US-09-030-607-32/c ^~~mence 32, Application US/09030607

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ATTORNEY AGENT INFORMATION:

NAME: Maki, David J.

REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 2101:
TELECOMMUNICATION INFORMATION:
TELEPAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
LENGTH: 789 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
US-09-030-607-32
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GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/030,607
FILLING DATE: 25-FEB-1998
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Xu, Jiangchun C. APPLICANT: Dillon, Davin C. TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS NUMBER OF SEQUENCES: 224
CORRESPONDENCE ADDRESS:
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Pred. No. 4.2e-80;
0; Mismatches 154;
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GGTAANACCACTATTCCTAATNGGTGGTGAGGGGNTGCGACCGCATTGTGCCCGTGGAAA

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US-09-439-313-32/c
; Sequence 32, Application US/0943931
; Patent No. 6329505
; GENERAL INFORMATION:
; APPLICANT: Xu, Jiangchun
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Harlocker, Susan Louis
; APPLICANT: Jiang Yuqui
; APPLICANT: Read, Steven G.
; APPLICANT: Kalos, Michael
; APPLICANT: Kalos, Michael
; APPLICANT: Fanger, Gary
; APPLICANT: Fanger, Gary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Jiang Yuqui
APPLICANT: Reed, Steven G.
APPLICANT: Relos, Michael
APPLICANT: Ralos, Michael
APPLICANT: Retter, Gary
APPLICANT: Retter, Mark
APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CUURRENT APPLICATION NUMBER: US/09/439,313
CUURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOPTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
SEQ ID NO 32
LENGTH: 789
TYPE: DNA
OFGAINION LATER CARGE
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; FEATURE;
; NAME/KEY: misc_feature
; LOCATION: (1)...(789)
; OTHER INFORWATION: n = A
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                                                                               G----CCCATGACCTTCGGCCTGGCCTGCTGCGCGTGGAGATGATGCACATGGCAGC-A
                                                                                                                        GGATTTGGGGGCCAAGGGGGATGACTTGTTGAAACTGGGNCCCCCCGGAGNTTTTTTGN
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llarity 75.9%;
Conservative
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Pred. No. 4.2e-80;
0; Mismatches 154;
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GENEAUT: DIILU.,
APPLICANT: Harlocker, Susan ......
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OP PROSTATE CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121,42708
CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT APPLICATION NUMBER: US/09/352,616A
NUMBER OF SEQ ID NOS: 472
NUMBER OF SEQ ID NOS: 472
COFTWARE: FastSEQ for Windows Version 3.0
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US-09-352-616A-32/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 32, Application Patent No. 6395278 GENERAL INFORMATION:
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Best Local Similarity
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NAME/KEY: misc_feature
LOCATION: (1)...(789)

OTHER INFORMATION: n = A,T,C

-09-352-616A-32
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75.9%;
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Pred. No. 4.2e-80;
0; Mismatches 154; Indels 13
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GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun

APPLICANT: Mitcham, Davin C.

APPLICANT: Mitcham, Jennifer Lynn

TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE

TITLE OF INVENTION: CANCER AND METHODS FOR THEIR USE

FILE REFERENCE: 210121.427C6

CURRENT APPLICATION NUMBER: US/09/232,149A

CURRENT FILING DATE: 1999-01-15

NUMBER OF SEQ ID NOS: 338

SOFTWARE: FastSEQ for Windows Version 3.0

LENGTH: 789
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US-09-232-149A-32/c
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                                                                                                                                                                                                              LENGTH: 789

TYPE: DNA

ORGANISM: Homo sapien
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(789)
                                                                                                                                                                                                    OTHER INFORMATION:
201 CACCCAGCCTGCCTAAAGGCCAGAGCCGTGGCTCCCAAACCCAGCAGCAGCCGGGCGA
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                                                                              GGGCCCGGCTGTGCAGGCACGAGGTGTCCATCAGAGCGTGGCCAACCGATGGCCCAAGCAG 200
                                                   GGGGGATGGTGNNNAGGANGGNGTTTTCNTTAGAGNNGGGCCCACCGGGGGCCCGAAGGG
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                                                                                                                Score 384.6; DB 3;
Pred. No. 4.2e-80;
0; Mismatches 154;
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APPLICANT: W. Jiangchun
APPLICANT: Dillon, Davin C.
FITTLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF
FILE REFERENCE: 210121. 428C5
CURRENT APPLICATION NUMBER: US/09/159,812A
CURRENT APPLICATION NUMBER: US/09/159,812A
CURRENT FILING DATE: 1998-09-23
NUMBER OF SEQ ID NOS: 306
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
SEQ ID NO 32
FENTURE:
NAME/KEY: misc_feature
LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
US-09-159-812-32
                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 7
US-09-159-812-32/c
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                                            Conservative
                                                       46.7%;
75.9%;
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                                         Score 384.6; DB 4;
Pred. No. 4.2e-80;
0; Mismatches 154;
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US-09-636-215-32/c
; Sequence 32, Application
; Patent No. 6620922
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             APPLICANT: Wang, Aijun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.42717C17
                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                       INFORMATION:
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    APPLICATION NUMBER: US/09/636,215
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                                                                                                              Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas (
Carter, Darrick
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Dillon, Davin C.
Mitcham, Jennifer
                                                                                                                                                                                                     Henderson, Robe
Kalos, Michael
                                                                                                  Li, Samuel
                                                                                                                                                                                                                                Jiang, Yuqui
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                                                                                                                                                                                                                  Robert
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RESULT 9
US-09-685-166A-32/c
Sequence 32, Application US
Sequent No. 6630305
Patent No. 6630305
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin
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NUMBER OF SEQ ID NOS: 852

SOFTWARE: FastSEQ for Windows V

SEQ ID NO 32

LENGTH: 789

TYPE: DNA

ORGANISM: Homo sapien
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(789)

OTHER INFORMATION: n = A,T,C or
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Pred. No. 4.2e-80;
0; Mismatches 154;
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; ORGANISM: Homo sapien
; PEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(789)
; OTHER INFORMATION: n = A
US-09-685-166A-32
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APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C21
CURRENT APPLICATION UNMERS: US/09/685,166A
CURRENT FILING DATE: 2000-10-10
NUMBER OF SEQ ID NOS: 898
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
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         AGAGGAAGATCAAGCGGGAGCGGAGGCTGCAGATCTGGTACCGCAGGTAGCGCCGCCGCC
                                                                            TCTACATCCCAGGCTGCCCACCTACGGCCGAGGCCCTGCTCTACGGCATCCTGCAGCTGC
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Vedvick, Thomas
Carter, Darrick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fanger, Gary R. Retter, Marc W. Stolk, John A.
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Kalos, Michael
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Pred. No. 4.2e-80;
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GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun

APPLICANT: Dillon, Davin C.

APPLICANT: Dillon, Davin C.

TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND TITLE OF INVENTION: METHODS FOR THEIR USE

FILE REFERENCE: 210121.427C4

CURRENT APPLICATION NUMBER: US/09/115,453B

CURRENT FILING DATE: 1998-07-14

NUMBER OF SEQ ID NOS: 228

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 32

LENGTH: 789

TYPE: DNA

ORGANISM: Homo Bapien

FEATURE:

NAME/KEY: misc_feature

LOCATION: (1)...(789)

OTHER INFORMATION: n = A,T,C or G
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US-09-115-453-32/c
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Best Local Similarity
Matches 527; Conserva
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                                               GCTACTACCACTATTCCTACTCGGTGGTGAGGGGCTGCGACCGGATCGTGCCCGTGGACA 610
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TCTACATCCCAGGCTGCCCACCTACGGCCGAGGCCCTGCTCTACGGCATCCTGCAGCTGC
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Pred. No. 4.2e-80;
0; Mismatches 154;
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APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE
FILE REFERENCE: 210121.427D2
CURRENT APPLICATION NUMBER: US/09/688,489
CURRENT FILING DATE: 2000-10-13
NUMBER OF SEQ ID NOS: 338
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
NAMES/KEY: misc_feature
LOCATION: (1):..(789)
COTHER INFORMATION: n = A,T,C or G
US-09-688-489-32
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US-09-688-489-32/c
; Sequence 32, Application
; Patent No. 6664377
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Best Local Similarity
Matches 527; Conserv
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AAGACCAGATGCCGGAGCCGCGANAAGTGGTTTCCANGGGGAG-TGNGCCAACGGNGGNG
                                         ACGACCAGATGCCGGAGCCGCGCTACGTGGTCTCCATGGGGAGCTGCGCCAACGGAGGAG
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Pred. No. 4.2e-80;
0; Mismatches 154;
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APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C20
CURRENT APPLICATION NUMBER: US/09/679,426
CURRENT FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 895
SOFTWARER: TALLING DATE: 1000-10-02
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US-09-679-426-32/c
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LENGTH: 789
                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapien
PEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C of
US-09-679-426-32
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APPLICANT:
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Local Similarity 75.9%;
nes 527; Conservative
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                                          GGANCCANCTTGCCNNCCAAAGNNAGAGGCGTGGGTTCCCAAACCCCAGNAANCGGGGGGA
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               GTATGTGGTGGCCAAGCTGGATGACCTCGT---CAACTGGGCCCGGCCGGAGTTCTCTGTG
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Dillon, Davin C.
Mitcham, Jennifer L.
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Kalos, Michael D.
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Pred. No. 4.2e-80;
0; Mismatches 154;
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JAPPLICANT: Wang, Aijun

APPLICANT: Wang, Aijun

APPLICANT: Skeiky, Yasir A.W.

APPLICANT: Hepler, William

TITLE OF INVENTION: COMPOSITIONS AND METH.

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE

FILE REFERENCE: 210121.427C23

CURRENT APPLICATION NUMBER: US/09/759,143

CURRENT FILING DATE: 2001-01-12

NUMBER OF SEQ ID NOS: 934

SOFTWARE: FastSEQ for Window?

SEQ ID NO 32

LENGTH: 789

TYPE: DNA

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Kalos, Michael
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Mitcham, Jennifer L.
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; FEATURE:

; MAME/KEY: misc feature

; LOCATION: (1)...(789)

; OTHER INFORMATION: n =

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Patent No. 6.
GENERAL INFO
APPLICANT
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Best Local Similarity 75.9
Matches 527; Conservative
IL INFORMATION:
ICANT: XU, Jiangchun
ICANT: XU, Jiangchun
ICANT: Dillon, Davin C.
ICANT: Mitcham, Jennifer L.
ICANT: Harlocker, Susan L.
ICANT: Jiang, Yuqui
ICANT: Henderson, Robert A
ICANT: Kalos, Michael D.
ICANT: Fanger, Gary R.
ICANT: Retter, Marc W.
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APPLICANT: Stolk, John A.

APPLICANT: Day, Craig H.

APPLICANT: Vedvick, Thomas S.

APPLICANT: Carter, Darrick

APPLICANT: Li, Samuel

APPLICANT: Li, Samuel

APPLICANT: Skeiky, Yasir A.W.

APPLICANT: Hepler, William

APPLICANT: Hepler, William

TITLE OF INVENTION: COMPOSITIONS AND METHODS

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CP

FILE REFERENCE: 210121.42718C18

CURRENT APPLICATION NUMBER: US/09/651,236

CURRENT APPLICATION NUMBER: US/09/651,236

CURRENT APPLICATION SEQ ID NOS: 865

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 32

LENGTH: 789

TYPE: DNA

ORGANISM: Homo sapien

FEATURE:

NAME/KEY: misc_feature

LOCATION: (1) ... (789)

OTHER INFORMATION: n = A,T,C or G

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                  GCCGCCGCCGGAGCCTGTCGCCGTCCTGTCCCCAGCCTGCTTGTGTCCCCGTGAGGTTGTC
                                                        GGATTTGGGGGGCCAAGGGGGATGACTTTGTTGAAACTGGGNCCCCCCGGAGNTTTTTTGN
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GCCGCCGCGAGCCTGTTGCCGTCCTGTCCCCAGCCTGNTTGTGTCCCGNGAGGTTGTC
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Pred. No. 4.2e-80;
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APPLICANT: Homburger et al.
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION UNMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 14895
TYPE: DNA
ORGANISM: Drosophila melanogaster
US-09-270-767-14895
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US-09-270-767-14895
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Search completed: March
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 33.7%;
Best Local Similarity 67.9%;
Matches 402; Conservative
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                 22,
                    2005, 04:10:50
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Pred. No. 3.8e-55;
0; Mismatches 189;
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